	changed a file from non-ASCII to ASCII ENTER E CRF Processing Date:
	Changed the margins in cases where the sequence text was "wrapped" down to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of fi page numbers throughout text; ☐ other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
•	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Deleted <i>ending</i> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (errollue to a Patentin bug). Sequences corrected:
	Other:

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*Examin r: Th abov corrections must be communicated to th applicant in the first Office Action. DO NOT send a copy of this form.

rage ror /



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Input Set : A:\pto.amc

Output Set: N:\CRF3\01152002\J020695.raw

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1 <110> APPLICANT: KIM, Dong-Soo
             NAM, Yoon-Kwon
      2
             NOH, Jae-Koo
      5 <120> TITLE OF INVENTION: EXPRESSION VECTOR CONTAINING LECTIN GENE REGULATION SITE OF
MUD
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      8 <130> FILE REFERENCE: PPBA0566
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RAW SEQUENCE LISTING DATE: 01/15/2002 PATENT APPLICATION: US/10/020,695 TIME: 16:55:53

Input Set : A:\pto.amc

Output Set: N:\CRF3\01152002\J020695.raw

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/020,695

DATE: 01/15/2002
TIME: 16:55:54

Input Set : A:\pto.amc

Output Set: N:\CRF3\01152002\J020695.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:221 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:225 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:229 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:234 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 2 L:235 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:239 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:243 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:251 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:255 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:259 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:284 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:284 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2 L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2

OIPE

RAW SEQUENCE LISTING DATE: 01/10/2002 PATENT APPLICATION: US/10/020,695 TIME: 15:54:13

Input Set : A:\ES.txt

12 <170> SOFTWARE: KopatentIn 1.71

Output Set: N:\CRF3\01102002\J020695.raw

ERRORED SEQUENCES

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604 <212> TYPE: DNA
605 <213> ORGANISM: Artificial Sequence
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608 <223> OTHER INFORMATION: primer for detecting a growth hormone gene of a carp
611 <400> SEQUENCE: 18
612 acaacacctg caccagctgg ctg
23
E--> 615 (1)

VERIFICATION SUMMARY

. . .

DATE: 01/10/2002

PATENT APPLICATION: US/10/020,695

TIME: 15:54:14

Input Set : A:\ES.txt

Output Set: N:\CRF3\01102002\J020695.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:221 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:225 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:229 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:234 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 2
L:235 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:239 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:243 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:251 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:255 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:259 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:284 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:288 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2